

FIGURE 1

MADPKYADLPGIARNEPDVYETSDLPEDDQAEFDAFAQELEELTSTSVEHIIVNPNAAYD  
KFKDKRVGKGLDFSDRIGKTKRTGYESGEYEMLGEGLGVKETPOOKYORLLHEVO  
ELTTEVEKIKTTVKESATEEKLTPVLLAKQLAALKQQLVASHLEKLLGPDAAINLTD  
DGALAKRLLLQLEATKNSKGGSGGKTTGTPDSSLVTYELHSRPEQDKFSQAAKVAELE  
KRLTELETAVRCDQDAQNPLSAGLQGACLMETVELLQAKVSALDLAVLDQVEARLQSV  
LGKVNEIAKHKASVEDADTQSKVHQLYETIQRWSPIASTLPQRLVTIKQLHEQAMQ  
FGQLLTHLDTTQQMANSKDNNTLLTQVQTTMRENLATVEGNFASIDERMKKLGK

007834-02401

**FIGURE 2**

MADPKYADLPGIARNEPDVYETSDLPEDDQAEFDAEELSSTSVEHIIVNPNAAYDKFKD  
KRVGTKGLDFSDRIGKTKRTGYESGDYEMLGEGLG**VKETPQOKYORLLHEVQELTT**  
**EVEKIKTTVKESATEEKLTPVVLAKQLAALKQQLVASHLEKLLGPDAAINLADPDGA**  
LAKRLLQLEA

007826-024404

### FIGURE 3

1 ATGGCCGATC CCAAGTTCCA GAACCTACCG GGAATAGCTT ATGACCAGCC  
51 GGACGTGTAC GAAACTCCAG ATGACCCGGA GCTCGATACA TCCGACTACT  
101 ACGAAGAGGA GCCGGAGAAC GAAGCCATCG AGCGACTGCA CATCTCGCCG  
151 AGCGTCGCTC ACAAGCGCTT CAGCGGAGCA ACGGTCGAGG GGAGTGTGGA  
201 CTTACGGAT CGCATTGGAC GACGCATGTG CCGGGGTTAC GATACGCGCG  
251 GCTCCAGCGA CTACGAGCTG GTTGGCCAGG GCGAGAAGGA GACGCCGGTG  
301 CAGAAGTGCC AGCGCCTGCA GATCGAGATG AACGAGCTTC TGAACGAGGT  
351 GGCCGCCTTG CAGGTGGACC GCAAGGTAGC CGACGAGGAG AAGCAGTCGT  
401 ACGATGCGGT GGCCACGGTT ATCAGCACGG CCCGAAAGGT GCTGGAGTCG  
451 CTGAAGCTGG AGCAAGTGCT GGGCAAGGAG CAGACGCCTG GAAGTAAGCA  
501 GGTGAAAGCA CTCATTAGCC AGGTGGAGGA GTTCAAGCAG TCCGGCGTTC  
551 TCACAGCCAT ACCCACGCCT GGCACCGATC TGGCGGCCAC GGCCCGCGTA  
601 GCCAGTCTAG AGCAGCGAAT CTCGCAGCTG GAGAAGGTGC TGGGCGCTCA  
651 GCCGGACAAG TTGAGCCGCC TTACCGCCGC CACCAACACC ACCAATGTAC  
701 TAGAGGCAGT GCGTCATCTA AGCACCAAGG CGGCCCTGAT ACAGCCTGAT  
751 AAATGGACA CCATCGAGCA GCGCCTGACC TCGCTGGCCG GCAAGATGGA  
801 TGCTATCGCC GAAAAGTCCA GCGGCAGTGC CCAGGACGCC AAACGAGATC  
851 AGAAGATTAC GGAATATAC GACATCGCGA AGCGCACGGA GCCAGTGGTG  
901 GAAATACTGC CGCACGTCAT CGAACGCATG CAAGCCCTCG AGGCCCTCCA  
951 TAAATATGCA AACAATTTCTG CCAAGATCAT CGCAGAGATT GAGCAGAAGC  
1001 AGGGAACCAT CACCACTAGC TTGGTGAACA ACAAGGAGCT GCTGCATTCC  
1051 GTACAGGAGA CTTTCGCCCA GAATCTGGAG ACTATCAACA GCAAGGTGGC  
1101 CAAGGTGGAG CAGCGTGTGG CGGCCATATC GTCTGCCAAA TGA

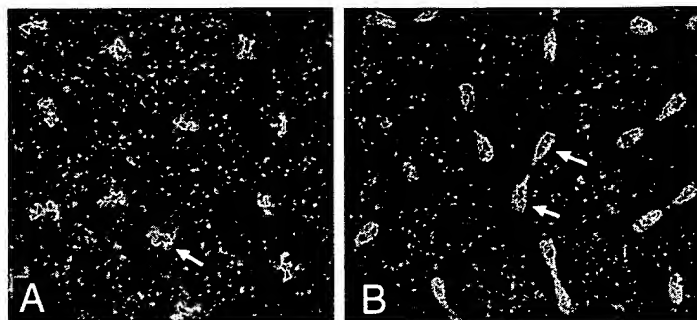
Figure 3: Sequence

**FIGURE 4**

MADPKFQNLPGIAYDQPDVYETPDDPELDTSDYYEEEPENEAIERLHISPSVAHKRFSGA  
TVEGSVDFTDRIGRRMCRGYDTRGSSDYELVGQGEKETPVQKCORLOIEMNELLNEV  
AALQVDRKVADEEKQSYDAVATVISTARKVLESCLKLEQVLGKEQTPGSKQVKALISQ  
VEEFKQSGVLTAIPTPGTDLAATARVASLEQRISQLEKVLGAQPDKLSRLTAATNTTNVL  
EAVRHLSTKAALIQPKLDTIEQRLTSLAGKMDAIAEKSSGSAQDAKRDQKITELYDIAK  
RTEPVVEILPHVIERMQALEALHKYANNFAKIIAEIEQKQGTITTSLVNNKELLHSVQETF  
AQNLETINSKVAKVEQRVAAISSAK

FIGURE 4

Control-injected



p50 peptide injected

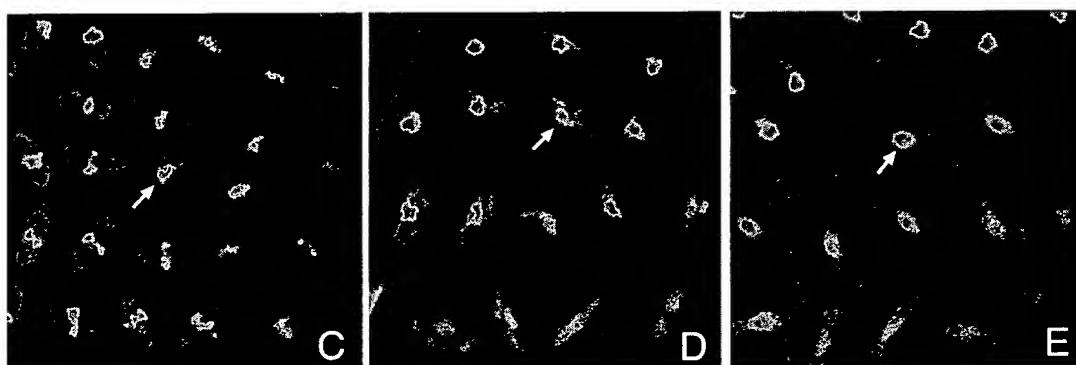


FIGURE 5